

AMENDMENTS TO THE CLAIMS

1.-80. (Canceled)

81. **(Currently amended)** A method for monitoring a response to a therapeutic protocol to treat prevent infection by a hepatitis virus pathogenic-agent, said method comprising determining the level of a cell surface marker selected from the group consisting of Toll-like receptor-2 (TLR-2) and Toll-like receptor-4 (TLR-4), ~~receptors and homologs thereof~~ wherein the efficacy of said therapeutic response is determined by a change in said level.

82. **(Previously presented)** The method of claim 81, wherein said level is compared to a sample selected from the group consisting of a pre-treatment sample and a control sample.

83. (Canceled)

84. **(Previously presented)** The method of claim 81, wherein said marker is affected in a manner selected from the group consisting of up-regulated and down-regulated.

85. **(Previously presented)** The method of claim 81, wherein said marker is determined by analyzing the mRNA or protein associated with said marker.

86. **(Currently amended)** The method of claim 81, wherein said hepatitis virus pathogenic-agent is selected from the group consisting of Salmonella, Escherichia, Klebsiella, Pasteurella, Bacillus, Clostridium, Corynebacterium, Mycoplasma, Ureaplasma, Actinomyces, Mycobacterium, Chlamydia, Chlamydomphila, Leptospira, Spirochaeta, Borrelia, Treponema, Pseudomonas, Burkholderia, Dichelobacter, Haemophilus, Ralstonia, Xanthomonas, Moraxella, Acinetobacter, Branhamella, Kingella, Erwinia, Enterobacter, Aerobacter, Citrobacter, Proteus, Providencia, Yersinia, Shigella, Edwardsiella, Vibrio, Rickettsia, Coxiella, Ehrlichia, Arcobacteria, Peptostreptococcus, Candida, Aspergillus, Trichomonas, Bacterioides, Coccidiomycetes, Pneumocystis, Cryptosporidium, Porphyromonas, Actinobacillus, Lactococcus, Lactobacillus, Zymomonas, Saccharomyces, Propionibacterium, Streptomyces, Penicillium, Neisseria, Staphylococcus, Campylobacter, Streptococcus, Enterococcus, Helicobacter, human immunodeficiency virus (HIV), Varicella-Zoster virus (VZV), herpes simplex virus (HSV), human papillomavirus (HPV), Hepatitis B virus (HBV), Hepatitis A virus (HAV), rhinovirus, echovirus, Coxsackievirus, cytomegalovirus, flavivirus, Ebola virus, paramyxovirus, influenza virus, enterovirus, Epstein-Barr virus, Marburg virus, polio virus, rabies virus, rubella virus, smallpox virus, rubeola virus, vaccinia virus, adenovirus, rotavirus and Hepatitis C virus (HCV) and Hepatitis B virus (HBV).

87. **(Currently amended)** A method for monitoring a response to a therapeutic protocol to treat prevent development of a disease condition, said disease condition resulting from infection by a hepatitis virus, said method comprising determining the level of a cell surface marker selected from the group consisting of TLR-2 and TLR-4, Toll-like receptors and homologs thereof wherein the efficacy of said therapeutic response is determined by a change in said level.

88. **(Previously presented)** The method of claim 87, wherein said level is compared to a sample selected from the group consisting of a pre-treatment sample and a control sample.

89. **(Canceled)**

90. **(Previously presented)** The method of claim 87, wherein said marker is affected in a manner selected from the group consisting of up-regulated and down-regulated.

91. **(Previously presented)** The method of claim 87, wherein said marker is determined by analyzing the mRNA or protein associated with said marker.

92. **(Currently amended)** The method of claim 87, wherein said hepatitis virus pathogenic agent is selected from the group consisting of Salmonella, Escherichia, Klebsiella, Pasteurella, Bacillus, Clostridium, Corynebacterium, Mycoplasma, Ureaplasma, Actinomyces, Mycobacterium, Chlamydia, Chlamydomphila, Leptospira, Spirochaeta, Borrelia, Treponema, Pseudomonas, Burkholderia, Dichelobacter, Haemophilus, Ralstonia, Xanthomonas, Moraxella, Acinetobacter, Branhamella, Kingella, Erwinia, Enterobacter, Arozoa, Citrobacter, Proteus, Providencia, Yersinia, Shigella, Edwardsiella, Vibrio, Rickettsia, Coxiella, Ehrlichia, Arcobacteria, Peptostreptococcus, Candida, Aspergillus, Trichomonas, Bacterioides, Coecidiomyces, Pneumocystis, Cryptosporidium, Porphyromonas, Actinobacillus, Lactococcus, Lactobacillus, Zymomonas, Saccharomyces, Propionibacterium, Streptomyces, Penicillium, Neisseria, Staphylococcus, Campylobacter, Streptococcus, Enterococcus, Helicobacter, human immunodeficiency virus (HIV), Varicella-Zoster virus (VZV), herpes simplex virus (HSV), human papillomavirus (HPV), Hepatitis B virus (HBV), Hepatitis A virus (HAV), rhinovirus, echovirus, Coxsackievirus, cytomegalovirus, flavivirus, Ebola virus, paramyxovirus, influenza virus, enterovirus, Epstein-Barr virus, Marburg virus, polio virus, rabies virus, rubella virus, smallpox virus, rubeola virus, vaccinia virus, adenovirus, rotavirus and Hepatitis C virus (HCV) and Hepatitis B virus (HBV).

93.-136. **(Canceled)**

137. **(Previously presented)** The method of Claim 81, wherein the change in the level of said cell surface marker is indicative of whether a subject will respond to a therapeutic intervention.

138 **(Previously presented)** The method of Claim 81, wherein the change in the level of said cell surface marker is predictive of an outcome of a therapeutic protocol.